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_			ttc Phe					-		-	_					307
_		_	act Thr	_					-	-					99	354
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Thr	Ala	Trp	Ala 20	Arg	Arg	Ser	Gln	Asp 25	Leu	His	Cys	Gly	A1a 30	Cys	Arg	
Ala	Leu	Va1 35	Asp	Glu	Leu	Glu	Trp 40		Ile	Ala	Gln	Va1 45		Pro	Lys	
Lys	Thr 50		Gln	Met	Gly	Ser 55	-	Arg	Пe	Asn	Pro 60	_	Gly	Ser	Gln	
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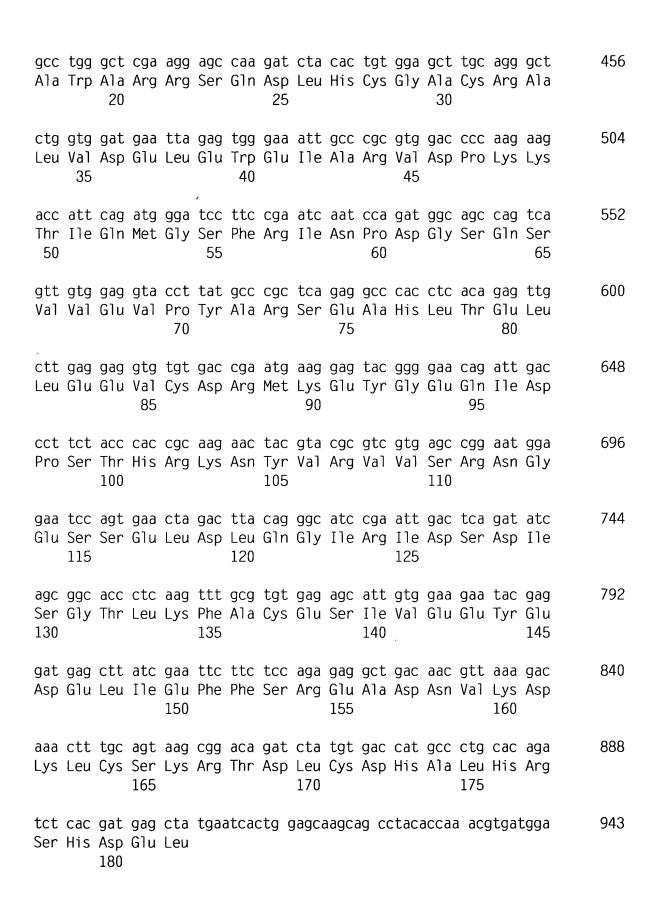
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		_	_	-			_		_	_		_	_	gtg Val 35	_	211
-		-		-		-	-		-		-	_		att Ile	•	259
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	cgc Arg															451
_	ctg Leu	_					•		_		•		_	~ ~		499
	aag Lys									-			-	-		547
	gaa Glu 150															595
	aag Lys			_		_	-		-	_			•		•	643
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Ala	Leu	Va1 35	Asp	Glu	Leu		Trp 40	Glu	Пe	Ala	Gln	Va1 45	Asp	Pro	Lys	

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                                                 125
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Glu Asp Glu Leu Ile Glu Phe Phe Ser Arg Glu Ala Asp Asn Val Lys
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cctgggccac ggcggcggcc gtgggagcag aggtggagcg accctgttac actaaag atg
                                                                      360
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                                                                 1
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1069

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Glu Asp Glu Leu Ile Glu Phe Phe Ser Arg Glu Ala Asp Asn Val Lys
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Asp Lys Leu Cys Ser Lys Arg Thr Asp Leu Cys Asp His Ala Leu His
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Arg Ser His Asp Glu Leu

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 Met Gly Ser Phe Arg Ile Asn Pro Asp Gly Ser Gln Ser Val Val Glu 35
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 Val Cys Asp Arg Met Lys Glu Tyr Gly Glu Gln Ile Asp Pro Ser Thr 65
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 75
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 His Arg Lys Asn Tyr Val Arg Val Val Ser Arg Asn Gly Glu Ser Ser 90
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 Glu Leu Asp Leu Gln Gly Ile Arg Ile Asp Ser Asp Ile Ser Gly Thr 100
 105
 110



Leu Lys	Phe 115	Ala	Cys	Glu	Ser	Ile 120	Val	Glu	Glu	Tyr	Glu 125	Asp	Glu	Leu
Ile Glu 130	Phe	Phe	Ser	Arg	Glu 135		Asp	Asn	Val	Lys 140		Lys	Leu	Cys
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